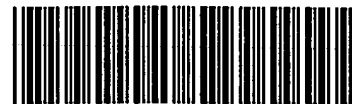


RAW SEQUENCE LISTING

**The Biotechnology Systems Branch of the Scientific and Technical
Information Center (STIC) no errors detected.**

Application Serial Number: 10/530,844A
Source: PCT
Date Processed by STIC: 3/10/06

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PCT

RAW SEQUENCE LISTING

DATE: 03/10/2006

PATENT APPLICATION: US/10/530,844A

TIME: 10:34:23

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Output Set: N:\CRF4\03092006\J530844A.raw

3 <110> APPLICANT: Consortium fur elektrochemische Industrie GmbH
 6 <120> TITLE OF INVENTION: Feedback-resistant homoserine-Transsuccinylases
 having a modified C-
 7 terminus
 10 <130> FILE REFERENCE: Co10221
 C--> 12 <140> CURRENT APPLICATION NUMBER: US/10/530,844A
 C--> 13 <141> CURRENT FILING DATE: 2005-04-08
 15 <160> NUMBER OF SEQ ID NOS: 12
 17 <170> SOFTWARE: PatentIn Ver. 2.0
 19 <210> SEQ ID NO: 1
 20 <211> LENGTH: 930
 21 <212> TYPE: DNA
 22 <213> ORGANISM: Escherichia coli
 24 <220> FEATURE:
 25 <221> NAME/KEY: CDS
 26 <222> LOCATION: (1)..(930)
 28 <300> PUBLICATION INFORMATION:
 29 <301> AUTHORS: Blattner, F. R.
 30 <302> TITLE: The complete genome sequence of Escherichia coli K-12.
 31 <303> JOURNAL: Science
 32 <304> VOLUME: 277
 33 <305> ISSUE: 533
 34 <306> PAGES: 1453-1474
 35 <307> DATE: 1997
 37 <400> SEQUENCE: 1
 38 atg ccg att cgt gtg ccg gac gag cta ccc gcc gtc aat ttc ttg cgt 48
 39 Met Pro Ile Arg Val Pro Asp Glu Leu Pro Ala Val Asn Phe Leu Arg
 40 1 5 10 15
 42 gaa gaa aac gtc ttt gtg atg aca act tct cgt gcg tct ggt cag gaa 96
 43 Glu Glu Asn Val Phe Val Met Thr Thr Ser Arg Ala Ser Gly Gln Glu
 44 20 25 30
 48 att cgt cca ctt aag gtt ctg atc ctt aac ctg atg ccg aag aag att 144
 49 Ile Arg Pro Leu Lys Val Leu Ile Leu Asn Leu Met Pro Lys Lys Ile
 50 35 40 45
 52 gaa act gaa aat cag ttt ctg cgc ctg ctt tca aac tca cct ttg cag 192
 53 Glu Thr Glu Asn Gln Phe Leu Arg Leu Leu Ser Asn Ser Pro Leu Gln
 54 50 55 60
 56 gtc gat att cag ctg ttg cgc atc gat tcc cgt gaa tcg cgc aac acg 240
 57 Val Asp Ile Gln Leu Leu Arg Ile Asp Ser Arg Glu Ser Arg Asn Thr
 58 65 70 75 80
 60 ccc gca gag cat ctg aac aac ttc tac tgt aac ttt gaa gat att cag 288
 61 Pro Ala Glu His Leu Asn Asn Phe Tyr Cys Asn Phe Glu Asp Ile Gln
 62 85 90 95
 64 gat cag aac ttt gac ggt ttg att gta act ggt gcg ccg ctg ggc ctg 336

RAW SEQUENCE LISTING

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TIME: 10:34:23

Input Set : A:\PTO.ss.txt

Output Set: N:\CRF4\03092006\J530844A.raw

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66              100              105              110
68  gtg gag ttt aat gat gtc gct tac tgg ccg cag atc aaa cag gtg ctg   384
69  Val Glu Phe Asn Asp Val Ala Tyr Trp Pro Gln Ile Lys Gln Val Leu
70              115              120              125
72  gag tgg tcg aaa gat cac gtc acc tcg acg ctg ttt gtc tgc tgg gcg   432
73  Glu Trp Ser Lys Asp His Val Thr Ser Thr Leu Phe Val Cys Trp Ala
74              130              135              140
76  gta cag gcc gcg ctc aat atc ctc tac ggc att cct aag caa act cgc   480
77  Val Gln Ala Ala Leu Asn Ile Leu Tyr Gly Ile Pro Lys Gln Thr Arg
78  145              150              155              160
80  acc gaa aaa ctc tct ggc gtt tac gag cat cat att ctc cat cct cat   528
81  Thr Glu Lys Leu Ser Gly Val Tyr Glu His His Ile Leu His Pro His
82              165              170              175
84  gcg ctt ctg acg cgt ggc ttt gat gat tca ttc ctg gca ccg cat tcg   576
85  Ala Leu Leu Thr Arg Gly Phe Asp Asp Ser Phe Leu Ala Pro His Ser
86              180              185              190
88  cgc tat gct gac ttt ccg gca gcg ttg att cgt gat tac acc gat ctg   624
89  Arg Tyr Ala Asp Phe Pro Ala Ala Leu Ile Arg Asp Tyr Thr Asp Leu
90              195              200              205
92  gaa att ctg gca gag acg gaa gaa ggg gat gca tat ctg ttt gcc agt   672
95  Glu Ile Leu Ala Glu Thr Glu Glu Gly Asp Ala Tyr Leu Phe Ala Ser
96              210              215              220
98  aaa gat aag cgc att gcc ttt gtg acg ggc cat ccc gaa tat gat gcg   720
99  Lys Asp Lys Arg Ile Ala Phe Val Thr Gly His Pro Glu Tyr Asp Ala
100  225              230              235              240
102  caa acg ctg gcg cag gaa ttt ttc cgc gat gtg gaa gcc gga cta gac   768
103  Gln Thr Leu Ala Gln Glu Phe Phe Arg Asp Val Glu Ala Gly Leu Asp
104              245              250              255
106  ccg gat gta ccg tat aac tat ttc ccg cac aat gat ccg caa aat aca   816
107  Pro Asp Val Pro Tyr Asn Tyr Phe Pro His Asn Asp Pro Gln Asn Thr
108              260              265              270
110  ccg cga gcg agc tgg cgt agt cac ggt aat tta ctg ttt acc aac tgg   864
111  Pro Arg Ala Ser Trp Arg Ser His Gly Asn Leu Leu Phe Thr Asn Trp
112              275              280              285
114  ctc aac tat tac gtc tac cag atc acg cca tac gat cta cgg cac atg   912
115  Leu Asn Tyr Tyr Val Tyr Gln Ile Thr Pro Tyr Asp Leu Arg His Met
116              290              295              300
118  aat cca acg ctg gat taa   930
119  Asn Pro Thr Leu Asp
120  305
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124 <211> LENGTH: 309
125 <212> TYPE: PRT
126 <213> ORGANISM: Escherichia coli
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130      1              5              10              15
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RAW SEQUENCE LISTING

DATE: 03/10/2006

PATENT APPLICATION: US/10/530,844A

TIME: 10:34:23

Input Set : A:\PTO.ss.txt

Output Set : N:\CRF4\03092006\J530844A.raw

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133           20           25           30
135 Ile Arg Pro Leu Lys Val Leu Ile Leu Asn Leu Met Pro Lys Lys Ile
136           35           40           45
138 Glu Thr Glu Asn Gln Phe Leu Arg Leu Leu Ser Asn Ser Pro Leu Gln
139           50           55           60
143 Val Asp Ile Gln Leu Leu Arg Ile Asp Ser Arg Glu Ser Arg Asn Thr
144           65           70           75           80
146 Pro Ala Glu His Leu Asn Asn Phe Tyr Cys Asn Phe Glu Asp Ile Gln
147           85           90           95
149 Asp Gln Asn Phe Asp Gly Leu Ile Val Thr Gly Ala Pro Leu Gly Leu
150           100          105          110
152 Val Glu Phe Asn Asp Val Ala Tyr Trp Pro Gln Ile Lys Gln Val Leu
153           115          120          125
155 Glu Trp Ser Lys Asp His Val Thr Ser Thr Leu Phe Val Cys Trp Ala
156           130          135          140
158 Val Gln Ala Ala Leu Asn Ile Leu Tyr Gly Ile Pro Lys Gln Thr Arg
159           145          150          155          160
161 Thr Glu Lys Leu Ser Gly Val Tyr Glu His His Ile Leu His Pro His
162           165          170          175
164 Ala Leu Leu Thr Arg Gly Phe Asp Asp Ser Phe Leu Ala Pro His Ser
165           180          185          190
167 Arg Tyr Ala Asp Phe Pro Ala Ala Leu Ile Arg Asp Tyr Thr Asp Leu
168           195          200          205
170 Glu Ile Leu Ala Glu Thr Glu Glu Gly Asp Ala Tyr Leu Phe Ala Ser
171           210          215          220
173 Lys Asp Lys Arg Ile Ala Phe Val Thr Gly His Pro Glu Tyr Asp Ala
174           225          230          235          240
176 Gln Thr Leu Ala Gln Glu Phe Phe Arg Asp Val Glu Ala Gly Leu Asp
177           245          250          255
179 Pro Asp Val Pro Tyr Asn Tyr Phe Pro His Asn Asp Pro Gln Asn Thr
180           260          265          270
182 Pro Arg Ala Ser Trp Arg Ser His Gly Asn Leu Leu Phe Thr Asn Trp
183           275          280          285
185 Leu Asn Tyr Tyr Val Tyr Gln Ile Thr Pro Tyr Asp Leu Arg His Met
186           290          295          300
190 Asn Pro Thr Leu Asp
191 305

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194 <210> SEQ ID NO: 3

195 <211> LENGTH: 34

196 <212> TYPE: DNA

197 <213> ORGANISM: Artificial Sequence

199 <220> FEATURE:

200 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer for

PCR

203 <400> SEQUENCE: 3

204 ctatttggtta gtgaataata gtactgagct ctgg

34

207 <210> SEQ ID NO: 4

208 <211> LENGTH: 34

209 <212> TYPE: DNA

210 <213> ORGANISM: Artificial Sequence

RAW SEQUENCE LISTING

DATE: 03/10/2006

PATENT APPLICATION: US/10/530,844A

TIME: 10:34:23

Input Set : A:\PTO.ss.txt

Output Set: N:\CRF4\03092006\J530844A.raw

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212 <220> FEATURE:
213 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer for
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216 <400> SEQUENCE: 4
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221 <211> LENGTH: 33
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223 <213> ORGANISM: Artificial Sequence
225 <220> FEATURE:
226 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer for
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229 <400> SEQUENCE: 5
230 tgggtggatat atgagatctg gtagacgtaa tag 33
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236 <211> LENGTH: 34
237 <212> TYPE: DNA
238 <213> ORGANISM: Artificial Sequence
240 <220> FEATURE:
241 <223> OTHER INFORMATION: Description of Artificial Sequence: Primer for
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244 <400> SEQUENCE: 6
245 gtatttgta gtgaataata gtactgagct ctgg 34
248 <210> SEQ ID NO: 7
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250 <212> TYPE: DNA
251 <213> ORGANISM: Artificial Sequence
253 <220> FEATURE:
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256 <400> SEQUENCE: 7
257 tcatatatcc accagctatt tgtagtgaa taa 33
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261 <211> LENGTH: 10
262 <212> TYPE: PRT
263 <213> ORGANISM: Artificial Sequence
265 <220> FEATURE:
266 <223> OTHER INFORMATION: Description of Artificial Sequence: Changes in
protein encoded by altered
267 Met A structural
268 gene
270 <400> SEQUENCE: 8
271 Ser Tyr Ile His Gln Leu Phe Val Ser Glu
272 1 5 10
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276 <211> LENGTH: 102
277 <212> TYPE: DNA
278 <213> ORGANISM: Artificial Sequence
280 <220> FEATURE:
281 <223> OTHER INFORMATION: Description of Artificial Sequence: Changes in
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285 <400> SEQUENCE: 9
286 tcatatatcc accactattt gtagtgaa aatagtactg agctctggat gcatacgct 60

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288 ttaattaagc ggccgcactg cgatgagtgg cagggcgggg cg

102

291 <210> SEQ ID NO: 10

RAW SEQUENCE LISTING

PATENT APPLICATION: US/10/530,844A

DATE: 03/10/2006

TIME: 10:34:23

Input Set : A:\PTO.ss.txt

Output Set: N:\CRF4\03092006\J530844A.raw

292 <211> LENGTH: 34

293 <212> TYPE: PRT

294 <213> ORGANISM: Artificial Sequence

296 <220> FEATURE:

297 <223> OTHER INFORMATION: Description of Artificial Sequence: Changes in Protein encoded by altered

298 Met A structural

299 gene

301 <400> SEQUENCE: 10

302 Ser Tyr Ile His His Tyr Leu Leu Val Asn Asn Ser Thr Glu Leu Trp

303 1 5 10 15

305 Met His Thr Arg Leu Ile Lys Arg Pro His Cys Asp Glu Trp Gln Gly

306 20 25 30

308 Gly Ala

312 <210> SEQ ID NO: 11

313 <211> LENGTH: 102

314 <212> TYPE: DNA

315 <213> ORGANISM: Artificial Sequence

317 <220> FEATURE:

318 <223> OTHER INFORMATION: Description of Artificial Sequence: Changes in Met A structural gene

320 <400> SEQUENCE: 11

321 tcatatatcc accagtatttt gttagtgaat aatagtactg agctctggat gcatacgcggt 60

323 ttaattaagc ggccgcactg cgatgagtgg cagggcgggg cg 102

326 <210> SEQ ID NO: 12

327 <211> LENGTH: 34

328 <212> TYPE: PRT

329 <213> ORGANISM: Artificial Sequence

333 <220> FEATURE:

334 <223> OTHER INFORMATION: Description of Artificial Sequence: Changes in protein encoded by altered

335 Met A structural

336 gene

338 <400> SEQUENCE: 12

339 Ser Tyr Ile His Gln Tyr Leu Leu Val Asn Asn Ser Thr Glu Leu Trp

340 1 5 10 15

342 Met His Thr Arg Leu Ile Lys Arg Pro His Cys Asp Glu Trp Gln Gly

343 20 25 30

345 Gly Ala

RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/530,844ADATE: 03/10/2006
TIME: 10:34:24

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VERIFICATION SUMMARY

DATE: 03/10/2006

PATENT APPLICATION: US/10/530,844A

TIME: 10:34:24

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L:13 M:271 C: Current Filing Date differs, Replaced Current Filing Date